

Clinical Potential of C- and P-MRS

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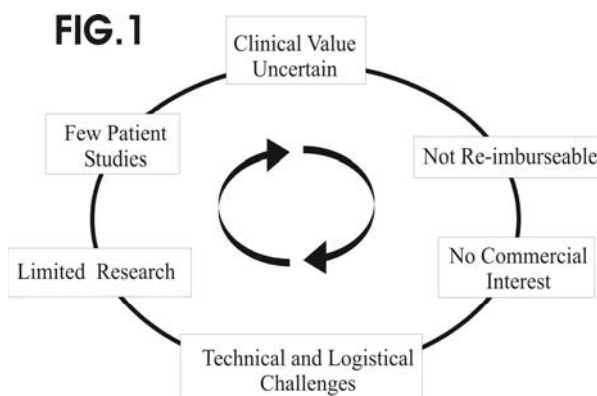
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Introduction

Proton MRS is by far the most widely used *in vivo* spectroscopy technique in humans. This is due to the fact that standard MRI hardware components are used, making ^1H MRS widely available, that the concentrations of protons are high, and that the MR sensitivity of protons is higher than the sensitivity of other nuclei. On the other hand proton MRS is sometimes limited by its lack of specificity. For example, elevated total choline in ^1H MRS could be due to an increase of phosphocholine (PC), glycerophosphocholine (GPC), or free choline or any combination. Cr comprises both free creatine and phosphocreatine. ^1H MRS is also compromised by low spectral resolution and complex pattern of some metabolites due to homonuclear J-coupling. Finally, important metabolites such as ATP are not observed with ^1H MRS.

Challenges for ^{13}C and ^{31}P MRS

Apart from identifying relevant medical and biological questions, there are significant challenges for both ^{13}C and ^{31}P MRS. Both methods compete with MRI, the probably most powerful diagnostic imaging tool, for valuable time on expensive equipment. In addition, multinuclear spectroscopy of disease is challenging because of the need for the availability of these tools on clinical MR scanners. This requires additional hardware, including dual-tuned RF coils and proton-decoupling^a capabilities which are not only expensive but sometimes, even if funds are available, hard to get installed on a clinical system. Both methods are also compromised by long acquisition times/low spatial resolution scans (due to low sensitivity)^b. Finally, in particular for ^{13}C MRS, expertise with data processing which includes complex mathematical modeling is needed.



Not surprisingly multinuclear MRS finds itself in the difficult situation illustrated in **Fig. 1**. However, as discussed below, both modalities offer unique opportunities to examine tissue *in vivo*.

^aProton-decoupling and the nuclear Overhauser effect (NOE) are essential in particular for direct detected ^{13}C MRS but also improve ^{31}P MRS considerably. It is not advised to attempt ^{13}C MRS without decoupling/NOE. Even the simplest application, such as measuring the lipid profile of a human leg would be quite demanding without the improved SNR and resolution facilitated by decoupling and NOE.

^bAn exception is ^{31}P MRS of skeletal muscle (e.g. leg). Due to the proximity of a surface coil to the leg muscle spectra of excellent quality can be obtained within a few seconds.

¹³C MRS

Only a few groups have attempted ¹³C MRS *in vivo*. The few studies undertaken illustrate the great promise of *in vivo* ¹³C MRS and interest in ¹³C MRS has grown considerably in recent years. A recent issue of *NMR in Biomedicine* (1) was exclusively dedicated to the application of ¹³C MRS to study biological systems. The potential of ¹³C arises from its biggest handicap: Low natural abundance of ¹³C (¹³C ≈ 1.1%, ¹²C ≈ 98.9%) and the compromised sensitivity (≈ 1/50 of ¹H) renders ¹³C MR spectroscopy *in vivo* very difficult due to the inherently very low signal-to-noise ratio (SNR). This low natural abundance, on the other hand, is the key to new, exciting applications of *in vivo* MRS:

¹³C MRS enables the investigation of metabolic pathways and the measurement of flux rates *in vivo* in humans after ¹³C enriched substrate infusion.

Fig. 2 illustrates the steps involved for a typical ¹³C MRS study after substrate infusion. *In vivo* ¹³C studies of humans require large amounts of ¹³C enriched substrates (expensive!), are technically and logistically challenging, and thus need to be planned carefully (2-4). Depending on the biological question asked, it needs to be decided what substrate should be infused for how long and in what fashion. For some applications oral administration may be appropriate (5-7) which would simplify the procedure

considerably because one intravenous (i.v.) infusion line could be eliminated.

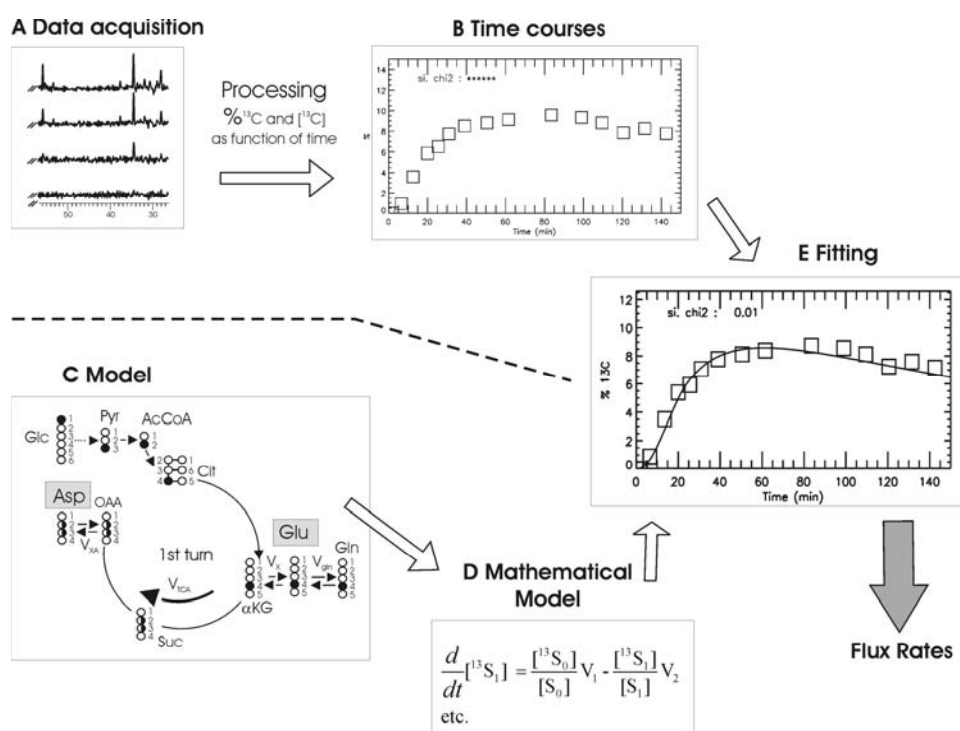


FIG. 2. (A) Dynamic ¹³C MRS involves the sequential acquisition of ¹³C spectra (typically every 2 – 10 minutes) after substrate infusion to monitor ¹³C label accumulation. (B) The next step is the quantitation of spectra to generate time courses of ¹³C concentration (or ¹³C enrichment). (C+D) A set of differential equations is derived from the metabolic model. (E) By iteratively varying flux parameters (and pool concentrations) of the mathematical model to optimize the fit with experimental data, flux rates (and pool concentrations) can be determined.

¹³C enriched glucose has been used by several groups to study brain metabolism *in vivo*. Glucose is the principal substrate for energy metabolism for both neurons and glia cells in the brain and facilitates the *de novo* synthesis of many neurochemicals. Glucose is the first choice as substrate because of its rapid oxidation and the fast appearance of ¹³C label in its metabolites and

its non-toxicity even at extremely high concentrations. Intravenously infused or orally administered glucose passes the blood-brain barrier and is readily metabolized through glycolysis and complete oxidation in the tricarboxylic acid (TCA)-cycle. ^{13}C enrichment of individual carbon atoms of glutamate (Glu), glutamine (Gln), aspartate (Asp), N-acetylaspartate (NAA), γ -amino butyric acid (GABA), lactate (Lac), alanine (Ala), and bicarbonate (HCO_3^-) follows and the *in vivo* rates of several of the principal bioenergetic pathways of the brain have been determined (2,3,8-11).

Applications

Potential clinical applications include diseases associated with physiological and pathological alterations in cerebral glucose consumption. An example is hepatic encephalopathy where cerebral glucose metabolism is altered, in response to hyperammonemia (12,13). Indeed, striking abnormalities in glucose metabolism and glutamate and glutamine label accumulation were observed in patients with chronic hepatic encephalopathy which appeared to be more pronounced with increasing severity of the disease (**Fig. 3**) (11). Inherited mitochondrial diseases of the

TCA-cycle or the electron transport chain also affect the overall rate of glucose oxidative metabolism in the brain and could be a target for ^{13}C MRS with glucose infusion. In addition to being the primary fuel of respiration, glucose is the source of α -glycerophosphate which is essential for biosynthesis of myelin phospholipids. ^{13}C MRS after ^{13}C enriched glucose infusion could therefore possibly contribute to understanding disorders in the developing brain including leukodystrophies and peroxisomal diseases which present as myelination defects on MRI. The information obtained from following the fate of ^{13}C labeled glucose goes beyond that of providing a rate for energy production. A tight coupling between cerebral glucose metabolism and glutamate neurotransmitter flux in humans has been proposed by Magistretti et al. (14). Aspartate, (a neurotransmitter?), can be studied *in vivo* in humans by its ^{13}C label accumulation. The role of NAA in mammalian brain, a neuronal/axonal marker which is central for its diagnostic power in ^1H MRS, is incompletely understood. NAA synthesis can be measured with ^{13}C MRS after glucose infusion in a clinical setting (5).

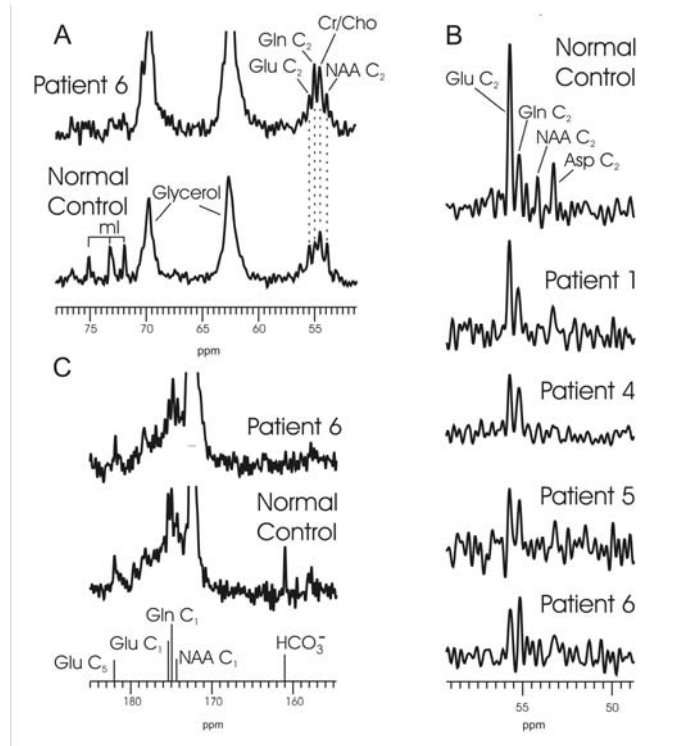


FIG. 3: *a:* Natural-abundance ^{13}C MRS of a control and severe chronic hepatic encephalopathy (CHE). Myo-inositol (mI) was depleted, glutamine C_2 resonance was strikingly elevated (threefold) and glutamate was modestly reduced (20%). *b:* Difference spectra from a control and four patients representing increasing stages of CHE, calculated from acquisitions 60-100 min after start of infusion. A progressive reduction of glutamate C_2 labeling was observed. *c:* ^{13}C spectra from the carbonyl region. At 60-100 min after start of infusion, ^{13}C incorporation into bicarbonate (the end product of complete glucose oxidation) is apparent in normal brain. In a severe case of grade III-IV CHE, bicarbonate enrichment was undetectable.

^{13}C labeled acetate: ^{13}C MRS studies employing ^{13}C enriched glucose as the substrate reflect predominantly the metabolism of neurons + glia. Acetate (Ac), an alternative fuel metabolized to acetyl-CoA only in the glial compartment (15) has been used in combination with ^{13}C glucose to separate glial from neuronal metabolism in cell preparations, tissue slices and *in vivo* in experimental animals (16,17). ^{13}C MRS after ^{13}C enriched acetate infusion therefore has the potential to investigate normal human glial metabolism and to elucidate brain diseases which originate in glia using the same MR technique as for glucose. Recent studies showed that the rate of Ac oxidation in human brain is $\approx 20\%$ of the total neuronal/glial TCA-cycle rate in fasted human brain (18,19). In patients with epileptic seizures on ketogenic diet abnormal label accumulation in glutamate and glutamine was observed (20) (**Fig. 4**). These results are consistent with altered glutamate-glutamine neurotransmitter cycling and adaptation to ketogenic diet with up-regulation of acetate oxidation relative to glucose oxidation.

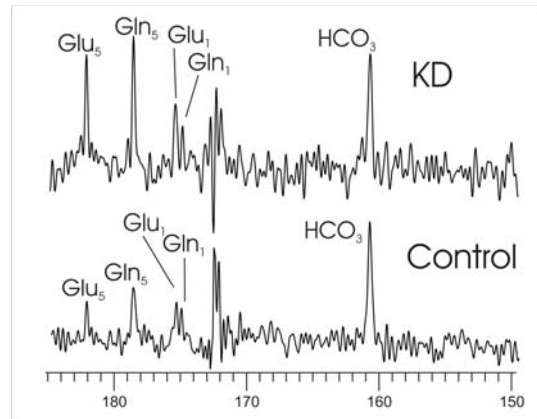
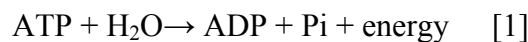


FIG. 4: Impact of ketogenic diet (KD) on astroglial acetate oxidation

(A) ^{13}C difference spectra obtained from ketogenic diet patients (average of three) and (B) controls after infusion of $[1-^{13}\text{C}]$ acetate. ^{13}C label incorporation into Glu C₅ and Gln C₅ was more pronounced in patients. In addition, Glu C₁ and Gln C₁ enrichment was more prominent, whereas equivalent production of HCO_3 was observed.

^{31}P MRS

Spectroscopy of living tissue started with ^{31}P MRS and it was not until technical advances in the early 1990s allowed efficient water suppression and reduction of eddy current artifacts that proton MRS become the dominant tool for *in vivo* spectroscopy. ^{31}P MRS allows the detection of phosphocreatine (PCr), ATP, and inorganic phosphate (Pi), and the measurement of pH from the chemical shift difference between PCr and Pi. ATP is the central provider of energy for all energy demanding processes in cells via the reaction:



PCr functions as a battery and is used to maintain/replenish ATP pools via **creatine kinase** when demand exceeds mitochondrial ATP production during strenuous exercise:

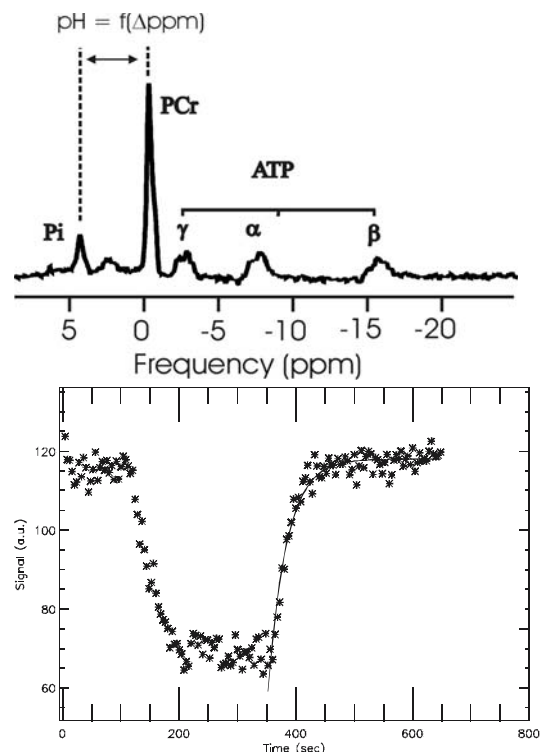
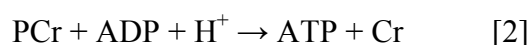
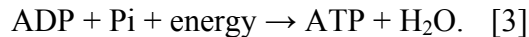


FIG. 5: ^{31}P MRS of skeletal muscle (top). Depletion and recovery of the muscle PCr levels during and after exercise (bottom).

Finally, at rest ATP is restored via oxidative phosphorylation:



ATP can then be used to replenish PCr via creatine kinase:



While creatine kinase requires the uptake of one proton, anaerobic glycolysis generates a proton from lactate acid. From the simultaneous measurements of pH and PCr important information about glycolysis and creatine kinase can be obtained. PCr/Pi is an indicator for adequate mitochondrial oxidative phosphorylation. The resting ^{31}P spectrum of muscle, even in disease, is often unremarkable. Some form of exercise, while continuously scanning, is required to study energy metabolism. In particular **skeletal muscle** is readily accessible and can be studied rested and during exercise with a time resolution of a few seconds (**Fig. 5**). This makes ^{31}P MRS a superb tool to study energy metabolism and mitochondrial function non-invasively in skeletal muscle and to diagnose and quantify the extent of metabolic myopathies.

^{31}P MRS of the diseased brain: A wide range of disease processes that include birth asphyxia, dementias, stroke, multiple sclerosis, epilepsy, mental disorders, hepatic encephalopathy, and tumors (21-39) were studied. The ^{31}P spectrum of **brain tissue** is more complex than spectra of muscle. In addition to ATP, PCr, and Pi, prominent phosphomonoester (PME) and phosphodiester (PDE) peaks are observed. **Proton-decoupling** is required to separate the PMEs phosphoethanolamine (PE), phosphocholine (PC) and the PDEs glycerophosphoethanolamine (GPE), glycerophosphocholine (GPC) from each other and from underlying broad signal from phospholipids (**Fig. 6**). These molecules are involved in myelin biosynthesis by methyl group metabolism and lipid transport and are components of a number of important biological compounds including the membrane phospholipids lecithin, sphingomyelin, and plasmalogen (40). Thus in the brain, ^{31}P MRS may provide useful information about (i) **energy metabolism** while it can be used at the same time to study diseases with (ii) **abnormal membrane metabolism**. In particular GPC and (possibly GPE) has been identified as a cerebral osmolyte (41,42) and ^{31}P MRS can be used to investigate disease which are believed to be associated with (iii) disturbances of **cerebral osmoregulation**. An example of the use of combined ^1H and

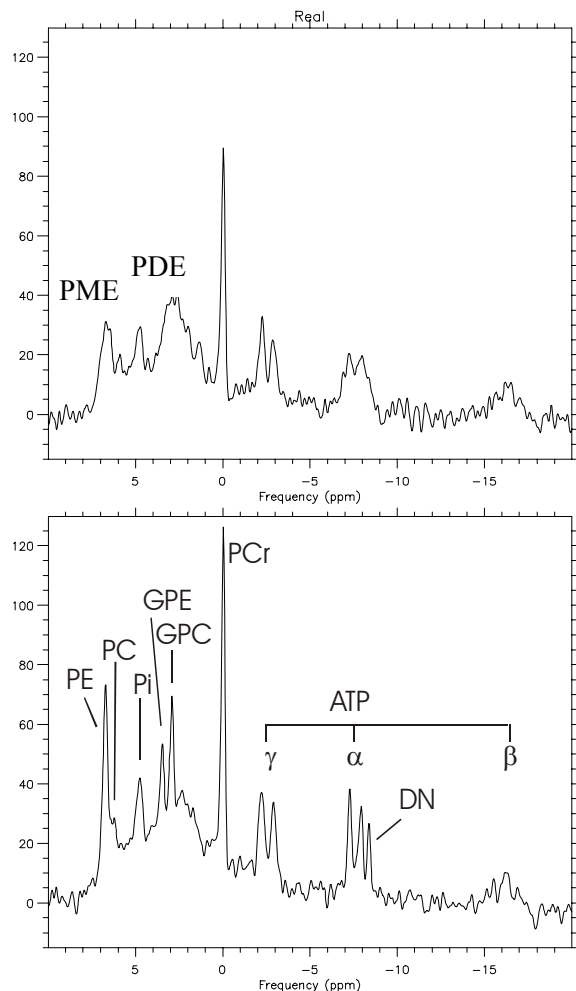
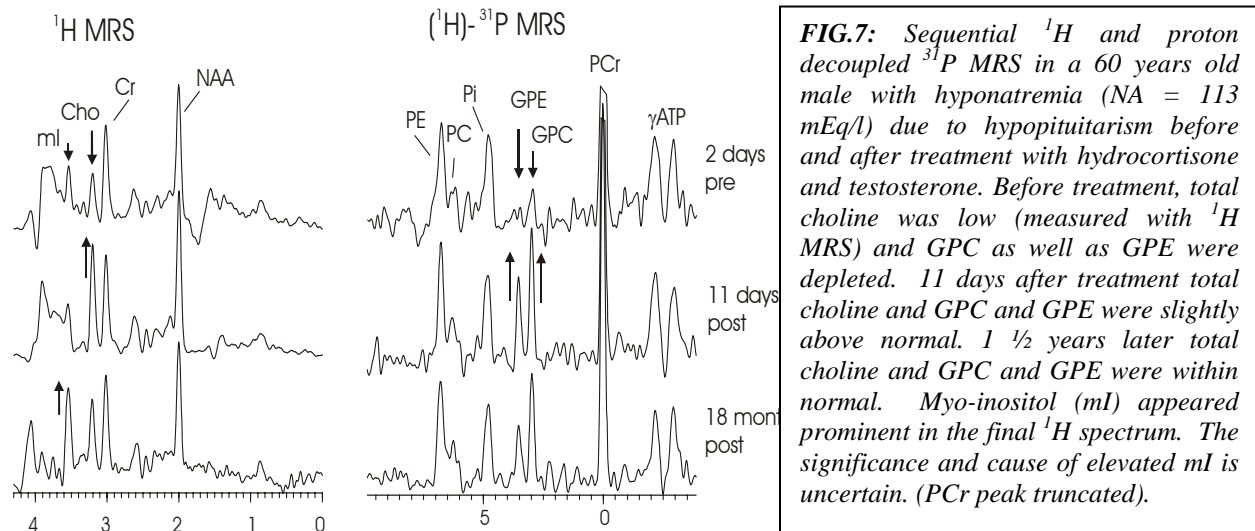


FIG. 6: ^{31}P (upper trace) and proton decoupled ^{31}P spectra of (normal) brain tissue. Proton decoupling is required to resolve peaks from PE, PC and GPE, GPC. By saturation of the protons a signal enhancement due to the Nuclear Overhauser Effect (NOE) is observed.

proton decoupled ^{31}P MRS to monitor the impact of treatment with hydrocortisone and testosterone of a subject with hypopituitarism is shown in **Fig. 7**.



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